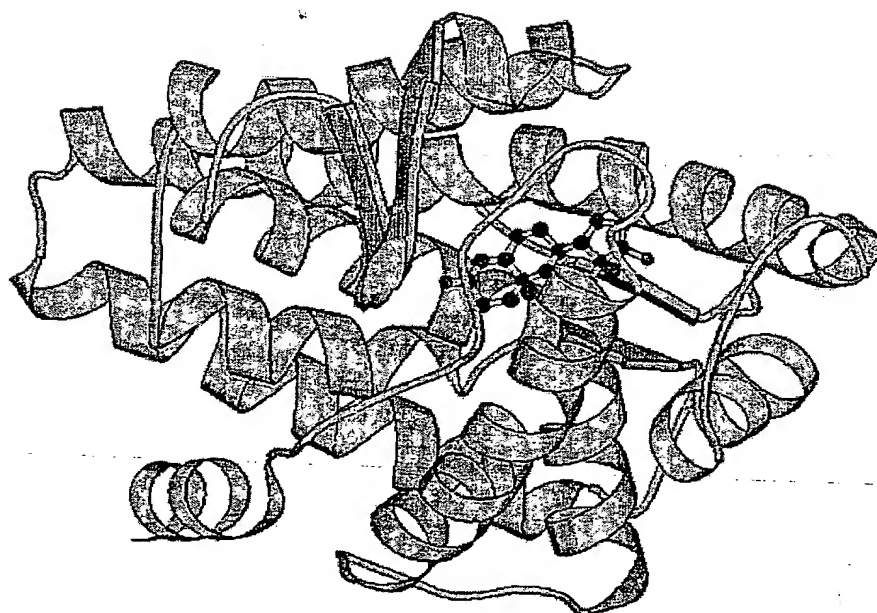


00667600 1000



**FIGURE 1**

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00692500-101200

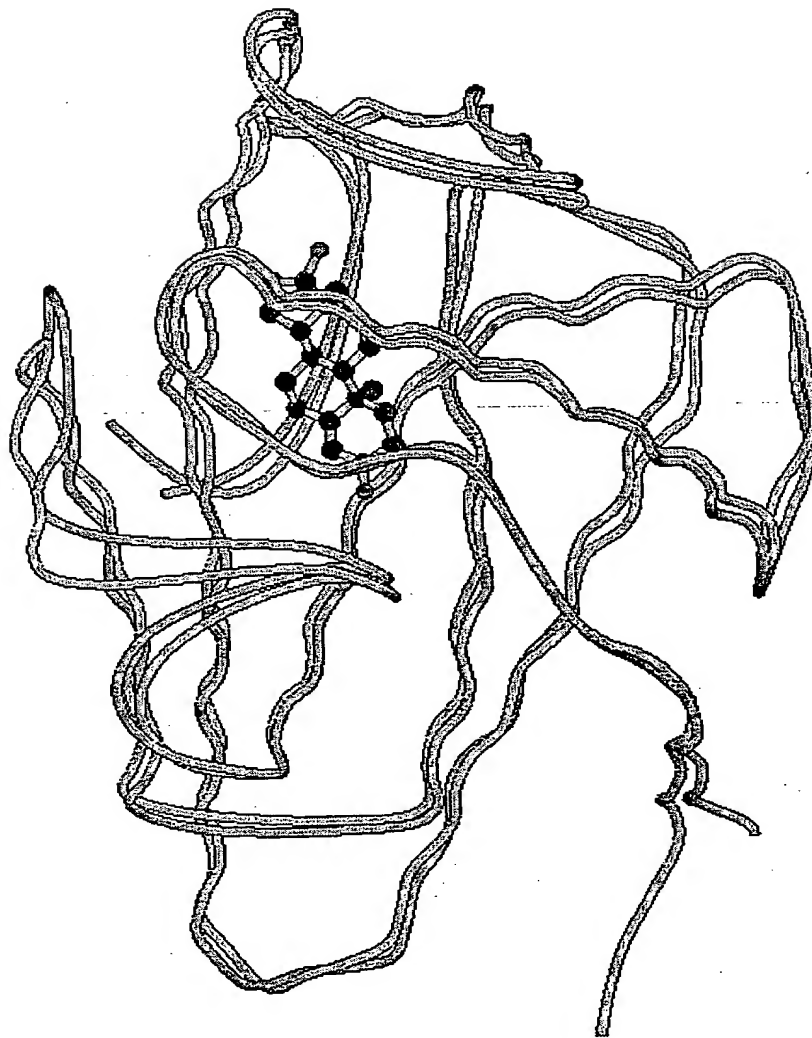


FIGURE 2

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00697600-101300

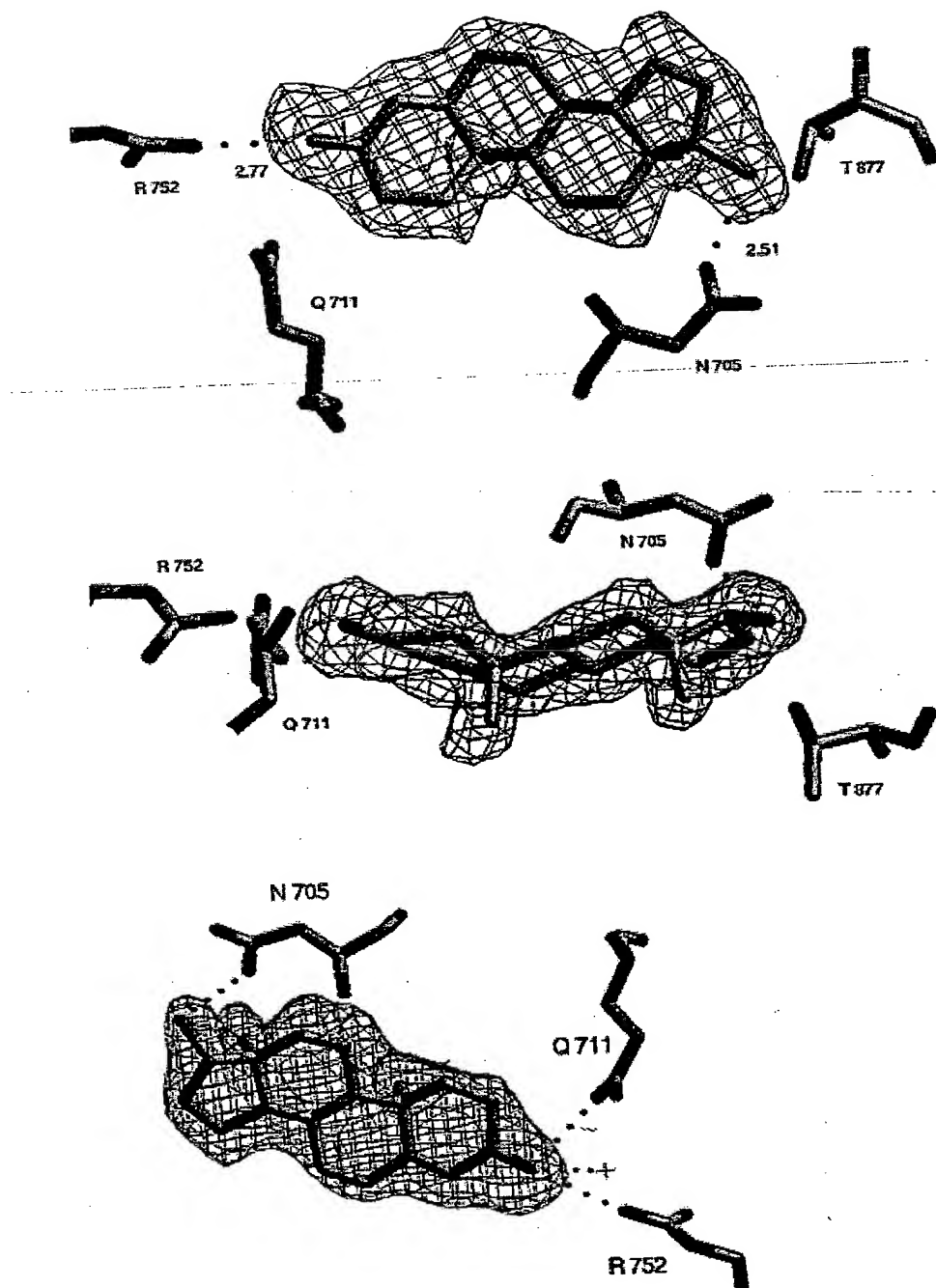


FIGURE 3

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The figure displays two panels of a molecular docking model, showing the interaction of a ligand (represented by a black stick model) with a protein structure (represented by a grey mesh). The ligand is a complex polycyclic molecule with a central aromatic core and several side chains. The protein structure is shown as a grey mesh, with specific residues highlighted in stick representation and labeled with their residue numbers. Dotted lines indicate hydrogen bonds between the ligand and the protein residues.

**Top Panel:** The ligand is docked in a binding pocket. Residues R752, Q711, N705, and T877 are shown in stick representation. Dotted lines indicate hydrogen bonds between the ligand and these residues.

**Bottom Panel:** The ligand is docked in a different binding pocket. Residues R766, Q725, N719, and C891 are shown in stick representation. Dotted lines indicate hydrogen bonds between the ligand and these residues.

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